



P6/09

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/831,088A

DATE: 08/26/2002
TIME: 14:36:51

Input Set : A:\pf0622usn_seqlist.txt
Output Set: N:\CRF3\08262002\I831088A.raw

4 <110> APPLICANT: INCYTE PHARMACEUTICALS, INC.
 5 TANG, Y. Tom
 6 CORLEY, Neil C.
 7 GUEGLER, Karl J.
 8 GORGONE, Gina A.
 9 AZIMZAI, Yalda
 10 KASER, Matthew R.
 11 YUE, Henry
 13 <120> TITLE OF INVENTION: COENZYME A-UTILIZING ENZYMES
 15 <130> FILE REFERENCE: PF-0622 PCT
 C--> 17 <140> CURRENT APPLICATION NUMBER: US/09/831,088A
 C--> 18 <141> CURRENT FILING DATE: 2002-08-26
 20 <150> PRIOR APPLICATION NUMBER: 09/185,217; unassigned
 W--> 21 <151> PRIOR FILING DATE: 1998-11-03; 1998-11-03
 23 <160> NUMBER OF SEQ ID NOS: 10
 25 <170> SOFTWARE: PERL Program
 27 <210> SEQ ID NO: 1
 28 <211> LENGTH: 100
 29 <212> TYPE: PRT
 30 <213> ORGANISM: Homo sapiens
 32 <220> FEATURE:
 33 <221> NAME/KEY: misc_feature
 34 <223> OTHER INFORMATION: Incyte ID No: 1580751CD1
 36 <400> SEQUENCE: 1
 37 Met Lys Gly Ala Thr Thr Asn Ile Cys Tyr Asn Val Leu Asp Arg
 38 1 5 10 15
 39 Asn Val His Glu Lys Lys Leu Gly Asp Lys Val Ala Phe Tyr Trp
 40 20 25 30
 41 Pro Cys Gln Arg Asp Gln Asp Gly Tyr Tyr Trp Ile Thr Gly Arg
 42 35 40 45
 43 Ile Asp Asp Met Leu Asn Val Ser Gly Glu Gly Gln Gly Pro Pro
 44 50 55 60
 45 Ser His Leu Ile Asn Ser Ala Pro Leu Thr Thr Pro Ser Arg Ser
 46 65 70 75
 47 Leu Pro Gln Glu Pro Arg Ser Val Leu Trp Pro Asp His Val Leu
 48 80 85 90
 49 Ser Val Ala Phe Ser Ser Gly Pro Arg Phe
 50 95 100
 54 <210> SEQ ID NO: 2
 55 <211> LENGTH: 159
 56 <212> TYPE: PRT
 57 <213> ORGANISM: Homo sapiens
 59 <220> FEATURE:

ENTERED

P6

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60 <221> NAME/KEY: misc_feature
61 <223> OTHER INFORMATION: Incyte ID No: 1627889CD1
63 <400> SEQUENCE: 2
64 Met Ser Gly Gln Gly Val Asp Arg His Leu Phe Ala Leu Tyr Ile
65 1 5 10 15
66 Val Ser Arg Phe Leu His Leu Gln Ser Pro Phe Leu Thr Gln Val
67 20 25 30
68 His Ser Glu Gln Trp Gln Leu Ser Thr Ser Gln Ile Pro Val Gln
69 35 40 45
70 Gln Met His Leu Phe Asp Val His Asn Tyr Pro Asp Tyr Val Ser
71 50 55 60
72 Ser Gly Gly Phe Gly Pro Ala Asp Asp His Gly Tyr Gly Val
73 65 70 75
74 Ser Tyr Ile Phe Met Gly Asp Gly Met Ile Thr Phe His Ile Ser
75 80 85 90
76 Ser Lys Lys Ser Ser Thr Lys Thr Asp Ser His Arg Leu Gly Gln
77 95 100 105
78 His Ile Glu Asp Ala Leu Leu Asp Val Ala Ser Leu Phe Gln Ala
79 110 115 120
80 Gly Gln His Phe Lys Arg Arg Phe Arg Gly Ser Gly Lys Glu Asn
81 125 130 135
82 Ser Arg His Arg Cys Gly Phe Leu Ser Arg Gln Thr Gly Ala Ser
83 140 145 150
84 Lys Ala Ser Met Thr Ser Thr Asp Phe
85 155
88 <210> SEQ ID NO: 3
89 <211> LENGTH: 215
90 <212> TYPE: PRT
91 <213> ORGANISM: Homo sapiens
93 <220> FEATURE:
94 <221> NAME/KEY: misc_feature
95 <223> OTHER INFORMATION: Incyte ID No: 1965888CD1
97 <400> SEQUENCE: 3
98 Met Asn Arg Pro Ser Ala Arg Asn Ala Leu Gly Asn Val Phe Val
99 1 5 10 15
100 Ser Glu Leu Leu Glu Thr Leu Ala Gln Leu Arg Glu Asp Arg Gln
101 20 25 30
102 Val Arg Val Leu Leu Phe Arg Ser Gly Val Lys Gly Val Phe Cys
103 35 40 45
104 Ala Gly Ala Asp Leu Lys Glu Arg Glu Gln Met Ser Glu Ala Glu
105 50 55 60
106 Val Gly Val Phe Val Gln Arg Leu Arg Gly Leu Met Asn Asp Ile
107 65 70 75
108 Ala Ser Ser Ala Val Met Gly Leu Ile Glu Thr Thr Arg Gly Leu
109 80 85 90
110 Leu Pro Gly Ala Gly Gly Thr Gln Arg Leu Pro Arg Cys Leu Gly
111 95 100 105
112 Val Ala Leu Ala Lys Glu Leu Ile Phe Thr Gly Arg Arg Leu Ser
113 110 115 120

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114 Gly Thr Glu Ala His Val Leu Gly Leu Val Asn His Ala Val Ala
 115 125 130 135
 116 Gln Asn Glu Glu Gly Asp Ala Ala Tyr Gln Arg Ala Arg Ala Leu
 117 140 145 150
 118 Ala Gln Glu Ile Leu Pro Gln Ala Pro Ile Ala Val Arg Leu Gly
 119 155 160 165
 120 Lys Val Ala Ile Asp Arg Gly Thr Glu Val Asp Ile Ala Ser Gly
 121 170 175 180
 122 Met Ala Ile Glu Gly Met Cys Tyr Ala Gln Asn Ile Pro Thr Arg
 123 185 190 195
 124 Asp Arg Leu Glu Gly Met Ala Ala Phe Arg Glu Lys Arg Thr Pro
 125 200 205 210
 126 Lys Phe Val Gly Lys
 127 215
 130 <210> SEQ ID NO: 4
 131 <211> LENGTH: 720
 132 <212> TYPE: PRT
 133 <213> ORGANISM: Homo sapiens
 135 <220> FEATURE:
 136 <221> NAME/KEY: misc_feature
 137 <223> OTHER INFORMATION: Incyte ID No: 2816341CD1
 139 <400> SEQUENCE: 4
 140 Met Asn Asn His Val Ser Ser Lys Pro Ser Thr Met Lys Leu Lys
 141 1 5 10 15
 142 His Thr Ile Asn Pro Ile Leu Leu Tyr Phe Ile His Phe Leu Ile
 143 20 25 30
 144 Ser Leu Tyr Thr Ile Leu Thr Tyr Ile Pro Phe Tyr Phe Phe Ser
 145 35 40 45
 146 Glu Ser Arg Gln Glu Lys Ser Asn Arg Ile Lys Ala Lys Pro Val
 147 50 55 60
 148 Asn Ser Lys Pro Asp Ser Ala Tyr Arg Ser Val Asn Ser Leu Asp
 149 65 70 75
 150 Gly Leu Ala Ser Val Leu Tyr Pro Gly Cys Asp Thr Leu Asp Lys
 151 80 85 90
 152 Val Phe Thr Tyr Ala Lys Asn Lys Phe Lys Asn Lys Arg Leu Leu
 153 95 100 105
 154 Gly Thr Arg Glu Val Leu Asn Glu Glu Asp Glu Val Gln Pro Asn
 155 110 115 120
 156 Gly Lys Ile Phe Lys Lys Val Ile Leu Gly Gln Tyr Asn Trp Leu
 157 125 130 135
 158 Ser Tyr Glu Asp Val Phe Val Arg Ala Phe Asn Phe Gly Asn Gly
 159 140 145 150
 160 Leu Gln Met Leu Gly Gln Lys Pro Lys Thr Asn Ile Ala Ile Phe
 161 155 160 165
 162 Cys Glu Thr Arg Ala Glu Trp Met Ile Ala Ala Gln Ala Cys Phe
 163 170 175 180
 164 Met Tyr Asn Phe Gln Leu Val Thr Leu Tyr Ala Thr Leu Gly Gly
 165 185 190 195
 166 Pro Ala Ile Val His Ala Leu Asn Glu Thr Glu Val Thr Asn Ile

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167	200	205	210
168	Ile Thr Ser Lys Glu Leu Leu Gln Thr	Lys Leu Lys Asp Ile Val	
169	215	220	225
170	Ser Leu Val Pro Arg Leu Arg His Ile	Ile Thr Val Asp Gly Lys	
171	230	235	240
172	Pro Pro Thr Trp Ser Glu Phe Pro Lys	Gly Ile Ile Val His Thr	
173	245	250	255
174	Met Ala Ala Val Glu Ala Leu Gly Ala	Lys Ala Ser Met Glu Asn	
175	260	265	270
176	Gln Pro His Ser Lys Pro Leu Pro Ser	Asp Ile Ala Val Ile Met	
177	275	280	285
178	Tyr Thr Ser Gly Ser Thr Gly Leu Pro	Lys Gly Val Met Ile Ser	
179	290	295	300
180	His Ser Asn Ile Ile Ala Gly Ile Thr	Gly Met Ala Glu Arg Ile	
181	305	310	315
182	Pro Glu Leu Gly Glu Glu Asp Val Tyr	Ile Gly Tyr Leu Pro Leu	
183	320	325	330
184	Ala His Val Leu Glu Leu Ser Ala Glu	Leu Val Cys Leu Ser His	
185	335	340	345
186	Gly Cys Arg Ile Gly Tyr Ser Ser Pro	Gln Thr Leu Ala Asp Gln	
187	350	355	360
188	Ser Ser Lys Ile Lys Lys Gly Ser Lys	Gly Asp Thr Ser Met Leu	
189	365	370	375
190	Lys Pro Thr Leu Met Ala Ala Val Pro	Glu Ile Met Asp Arg Ile	
191	380	385	390
192	Tyr Lys Asn Val Met Asn Lys Val Ser	Glu Met Ser Ser Phe Gln	
193	395	400	405
194	Arg Asn Leu Phe Ile Leu Ala Tyr Asn	Tyr Lys Met Glu Gln Ile	
195	410	415	420
196	Ser Lys Gly Arg Asn Thr Pro Leu Cys	Asp Ser Phe Val Phe Arg	
197	425	430	435
198	Lys Val Arg Ser Leu Leu Gly Gly Asn	Ile Arg Leu Leu Leu Cys	
199	440	445	450
200	Gly Gly Ala Pro Leu Ser Ala Thr Thr	Gln Arg Phe Met Asn Ile	
201	455	460	465
202	Cys Phe Cys Cys Pro Val Gly Gln Gly	Tyr Gly Leu Thr Glu Ser	
203	470	475	480
204	Ala Gly Ala Gly Thr Ile Ser Glu Val	Trp Asp Tyr Asn Thr Gly	
205	485	490	495
206	Arg Val Gly Ala Pro Leu Val Cys Cys	Glu Ile Lys Leu Lys Asn	
207	500	505	510
208	Trp Glu Glu Gly Gly Tyr Phe Asn Thr	Asp Lys Pro His Pro Arg	
209	515	520	525
210	Gly Glu Ile Leu Ile Gly Gly Gln Ser	Val Thr Met Gly Tyr Tyr	
211	530	535	540
212	Lys Asn Glu Ala Lys Thr Lys Ala Asp	Phe Phe Glu Asp Glu Asn	
213	545	550	555
214	Gly Gln Arg Trp Leu Cys Thr Gly Asp	Ile Gly Glu Phe Glu Pro	
215	560	565	570

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216 Asp Gly Cys Leu Lys Ile Ile Asp Arg Lys Lys Asp Leu Val Lys
 217 575 580 585
 218 Leu Gln Ala Gly Glu Tyr Val Ser Leu Gly Lys Val Glu Ala Ala
 219 590 595 600
 220 Leu Lys Asn Leu Pro Leu Val Asp Asn Ile Cys Ala Tyr Ala Asn
 221 605 610 615
 222 Ser Tyr His Ser Tyr Val Ile Gly Phe Val Val Pro Asn Gln Lys
 223 620 625 630
 224 Glu Leu Thr Glu Leu Ala Arg Lys Lys Gly Leu Lys Gly Thr Trp
 225 635 640 645
 226 Glu Glu Leu Cys Asn Ser Cys Glu Met Glu Asn Glu Leu Leu Lys
 227 650 655 660
 228 Val Leu Ser Glu Ala Ala Ile Ser Ala Ser Leu Glu Lys Phe Glu
 229 665 670 675
 230 Ile Leu Val Lys Ile Arg Leu Ser Pro Glu Pro Trp Thr Pro Glu
 231 680 685 690
 232 Thr Gly Leu Val Thr Asp Ala Phe Lys Leu Lys Arg Lys Glu Leu
 233 695 700 705
 234 Lys Thr His Tyr Gln Ala Asp Ile Glu Arg Met Tyr Gly Arg Lys
 235 710 715 720

238 <210> SEQ ID NO: 5

239 <211> LENGTH: 456

240 <212> TYPE: PRT

241 <213> ORGANISM: Homo sapiens

243 <220> FEATURE:

244 <221> NAME/KEY: misc_feature

245 <223> OTHER INFORMATION: Incyte ID No: 3324214CD1

247 <400> SEQUENCE: 5

248 Met Phe Leu Leu Leu Thr Phe Asp Ser Leu Ile Val Asn Leu Leu
 249 1 5 10 15
 250 Gly Ile Ser Leu Thr Val Leu Phe Thr Leu Leu Leu Val Phe Ile
 251 20 25 30
 252 Ile Val Pro Ala Ile Phe Gly Val Ser Phe Gly Ile Arg Lys Leu
 253 35 40 45
 254 Tyr Met Lys Ser Leu Leu Lys Ile Phe Ala Trp Ala Thr Leu Arg
 255 50 55 60
 256 Met Glu Arg Gly Ala Lys Glu Lys Asn His Gln Leu Tyr Lys Pro
 257 65 70 75
 258 Tyr Thr Asn Gly Ile Ile Ala Lys Asp Pro Thr Ser Leu Glu Glu
 259 80 85 90
 260 Glu Ile Lys Glu Ile Arg Arg Ser Gly Ser Ser Lys Ala Leu Asp
 261 95 100 105
 262 Asn Thr Pro Glu Phe Glu Leu Ser Asp Ile Phe Tyr Phe Cys Arg
 263 110 115 120
 264 Lys Gly Met Glu Thr Ile Met Asp Asp Glu Val Thr Lys Arg Phe
 265 125 130 135
 266 Ser Ala Glu Glu Leu Glu Ser Trp Asn Leu Leu Ser Arg Thr Asn
 267 140 145 150
 268 Tyr Asn Phe Gln Tyr Ile Ser Leu Arg Leu Thr Val Leu Trp Gly

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 08/26/2002
PATENT APPLICATION: US/09/831,088A TIME: 14:36:52

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#: 6; N Pos. 63

Seq#: 10; N Pos. 1597, 1650, 1656, 1713, 1719, 1754, 1778, 1791, 1794, 1796, 1800

VERIFICATION SUMMARY

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Input Set : A:\pf0622usn_seqlist.txt

Output Set: N:\CRF3\08262002\I831088A.raw

L:17 M:270 C: Current Application Number differs, Replaced Current Application Number
L:18 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:21 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:328 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:60
L:516 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:10
L:543 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:1560
L:544 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:1620
L:545 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:1680
L:546 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:1740